TOCABOLDO (1491, 1492) NOVAI FORMER MIN. ADBRING (1491, 1492) NOVAI FORMER MIN. ADBRING (1491, 1492) NOVAI FORMER MIN. ADBRING (1492, 1900) NOVAI FORM		10001 1001 11 10001 0001			
100417319 (2009, 2040) Nove Proteins am, Galank gil 193591 gipdi (Jodzea - ring finger protein 101322811 (1008, 1070) Nove Proteins am, Galank gil 1402657 (J00358) - cytoplasmic protein Nove Protein am, Galank gil 1402657 (J00358) - cytoplasmic protein Nove Protein am, Galank gil 1402657 (J00358) - cytoplasmic protein Nove Protein am, Galank gil 1402657 (J00358) - cytoplasmic protein Nove Protein am, Galank gil 1402657 (J00358) - cytoplasmic protein Nove Protein am, Galank gil 1402657 (J00358) - cytoplasmic protein Nove Protein am, Galank gil 1635699 cytoplasmic protein cytoplasmic c	382	100340205 (1481, 1482)	OLCA -	UNCLASSIFIED	1022, 1025, 1026, 1030, 1042
101322811 (1038, 1070) Nove Protein sin, GBank gill 402687 (U80598) - cyropisamic protein	383	100417319 (2039, 2040)		UNCLASSIFIED	1022, 1040, 1042
### 10132119 (\$65), \$64) Nove Protein Relay (g) 14302829 (m) Cologes ((277137) Myoch Protein Protein Relay (g) 146599 (m) Cologes ((277137) Myoch Protein Protein Relay (g) 1655999 (m) Cologes ((277137) Myoch Protein Relay (g) 167070 (m) Cologes ((277137) Myoch Protein Relay (g) 167070 (m) Cologes ((277137) Myoch Protein Relay (g) 167070 (m) Cologes ((277137) Myoch Protein Relay ((272137) Myo	384	101323811 (1069, 1070)	ein sim. GBank gi 1402857 (U60593) - cytoplasmic protein musculus]	UNCLASSIFIED	1022, 1037, 1040, 1042
10392094 (2049, 2050) Nove Protein sin. GBank gil (1655699 emb(CAA690022 - (Y07722) Accord Protein sin. GBank gil (15770 emb(CAA69022 - (Y07722) Accord Protein sin. GBank gil (17770 emb(CA69022 - (Y07256) Accord Protein sin. GBank gil (17770 emb(CA69022 - (Y02256) Accord Protein sin. GBank gil (17770 emb(CA69022 - (Y02256) Accord Protein sin. GBank gil (17770 emb(CA69022 - weak similarity to product (Caenomabditis elegans) Accord (Caenoma	385	87942019 (563, 564)	277137)	UNCLASSIFIED	1022
87928991 (345, 346) Novel Protein and Caleank gil1707706/psp[PG386504 (Y08250) 87919431 (947, 348) Novel Protein and Caleank gil1717706/psp[PG386517824_MOUSE - Protein and Caleank gil2706.CAFCO0036] - week similarity to Caleaners, gil2706.CAFCO00361 - week similarity to Caleancers gil2706.CAFCO000361 - week similarity to Caleancers gil2706.CAFCO000361 - week similarity to Caleancers gil27065.CAFCO000361 - week similarity to Caleance gil27065.CAFCO0000361 - week similarity to Caleance gil27065004 (U49973) - ORFT to Calean	386	101321949 (2049, 2050)	Y07752)	UNCLASSIFIED	1022, 1033
87919431 (947, 948) Novel Protein sim. Glaank gij 1777839ppPs39sgf7824_MOUSE - PROTEIN 1592_MOUSE - PROTEIN 15	387	87938991 (345, 346)	- (Y08256)	UNCLASSIFIED	1022
100399052 (1675, 1676) Nove Protein sim. GBank gil1938574 (U67180) - B0025.2 gene product (Caenomabditis elegans) 100399052 (1675, 1676) Nove Protein sim. GBank gil19347160 (AF000299) - weak similarity to collagens; glycine- and profilme-rich (Caenomabditis elegans) 87940066 (399, 400) Nove Protein sim. GBank gil2078493 (U4320) - mitteezal production sim. GBank gil2078493 (U4320) - mitteezal production sim. GBank gil2078493 (U4320) - mitteezal procedured mun tuberculosis 100344205 (1880, 1890) Nove Protein sim. GBank gil214473 (U96963) - pi 40mDia fMus musculusi	388	87919481 (947, 948)		UNCLASSIFIED	1022
100389052 (1675, 1676) Novel Protein sim. GBank gil 94760 (AF000289) - weak similarity to congagenes, glydree- and profuse-field (Caenorhabditis elegans)		100360209 (1891, 1892)		UNCLASSIFIED	1011, 1012, 1013,
100393052 (1675, 1676)			product [Caenorhabditis elegans]		1022,
100383052 (1675, 1676) Novel Protein sim. GBank gil1947160 (AF000298) - weak similarity to collaparies; glyche- and proline-rich (Caenorinabolite elegans) 87940065 (399, 400) Novel Protein sim. GBank gil2065310 GV27899 Movel Protein sim. GBank gil2065310 GV27899 Movel Protein sim. GBank gil207899 Movel Protein sim. GBank gil207899 Movel Protein sim. GBank gil20789 Movel Protein sim. GBank gil20789 GV3789 Movel Protein sim. GBank gil214473 (U99983) - p140mDla [Mus Musculus] 100344205 (1889, 1890) Novel Protein sim. GBank gil214473 (U99983) - p140mDla [Mus Musculus] 100344205 (1889, 1890) Novel Protein sim. GBank gil214473 (U99983) - p140mDla [Mus Musculus] 100393146 (1679, 1680) Novel Protein sim. GBank gil214470 [Ipil][IS5210 - tricarboxylate Cardiar - rat (Iraganse m) GBank gil214473 (U49973) - Graffer - rat (Iraganse m) GBank gil2139478 Movel Protein sim. GBank gil224054 [Ipil][IS5210 - tricarboxylate Cardiar - rat (Iraganse m) GBank gil2226005 (U49973) - ORF1; MER37; Novel Protein sim. GBank gil2226005 (U49973) - ORF1; MER37; Novel Protein sim. GBank gil2226005 (U49973) - ORF1; MER37; Novel Protein sim. GBank gil2226005 (U49973) - ORF1; turction Unknown [Homo sapiens] 87942813 (39, 40) Novel Protein sim. GBank gil22403pp O10341 V091_NPVOP - HYPOP - HY	389	-			1025, 1028, 1030, 1033, 1037, 1039,
87940065 (399, 400) Novel Protein sim. GBank gilgo65210[emb[GA473251] · (Y12713) 87917002 (411, 412) Novel Protein sim. GBank gilgo72269[emb[GA80333] · (255121) 87940483 (519, 520) Novel Protein sim. GBank gilgo72269[emb] or antifecace glycopeptide Argon (1900) - antifecace glycopeptide Glycopeptide (1900) - antifecace glycopeptide Glycopeptide (1900) - antifecace glycopeptide (1900) - antifeca	390	100393052 (1675, 1676)	ein sim. GBank gij1947160 (AF000298) - weak similarity to alvoine- and proline-rich [Caenorhabditis elegans]	UNCLASSIFIED	
### 87917002 (411, 412) Wover Protein sim. GBank gij2072699 emb CAB06339 - (295121)	391	87940065 (399, 400)		UNCLASSIFIED	1022
87940463 (519, 520) Novel Protein sim. GBank gi 2078483 (U43200) - antifreeze giveopeptide AFGP polyprotein precursor Boreogadus saidal 100344205 (1889, 1890) Novel Protein sim. GBank gi 2114473 (U96963) - p140mDia [Mus musculus] Novel Protein sim. GBank gi 2129478[pir][5S210 - tricarboxylate carrier - rat (fragment) and the protein sim. GBank gi 2144101[pir][5S210 - tricarboxylate carrier - rat (fragment) and the protein sim. GBank gi 214504 [1613, 1614) Novel Protein sim. GBank gi 2226004 (U49973) - ORF1; MER37; putative transposase similar to pogo element [Homo sapiens] Novel Protein sim. GBank gi 2226004 (U49973) - ORF2; tunction unknown [Homo sapiens] Novel Protein sim. GBank gi 2226005 [U49973) - ORF2; tunction unknown [Homo sapiens] Novel Protein sim. GBank gi 2226005 [U49973] - ORF2; tunction unknown [Homo sapiens] Novel Protein sim. GBank gi 2236240[sp](1975, 1976) Novel Protein sim. GBank gi 2436240[sp](10341 Y091_NVOP - HVPOTHETICAL 29.3 KD PROTEIN (ORF29) Novel Protein sim. GBank gi 243633[sp P75839]FLGG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN [FLGG (DISTAL ROD	392	87917002 (411, 412)		UNCLASSIFIED	1022
100349205 (1889, 1890) Novel Protein sim. GBank gil213473 (U96963) - p140mDla [Mus musculus] 100393146 (1679, 1680) Novel Protein sim. GBank gil2129478 pir SS1939 - chitinase (EC 3.2.1.14) precursor - beat 101334754 (1343, 1344) Novel Protein sim. GBank gil2144101 pir ISS210 - tricarboxylate carrier - rat (fragment) R7940670 (547, 548) Novel Protein sim. GBank gil215864 pir S72805 - B1549_F2_87 Novel Protein sim. GBank gil2228004 (U49973) - ORF1: MER37:	393	87940463 (519, 520)		UNCLASSIFIED	1022
100393146 (1679, 1680) Novel Protein sim. GBank gil2129478 pir S51939 - chitinase (EC 3.2.1.14) precursor - beet 101334754 (1343, 1344) Novel Protein sim. GBank gil214101 pir I55210 - tricarboxylate carrier - rat (fragment) R7940670 (547, 548) Novel Protein sim. GBank gil2245684 pir S72805 - B1549_F2_87 Rabidossis thallana R7931075 (1577, 1578) Novel Protein sim. GBank gil2226004 (U49973) - ORF1; MER37; putative transposaes similar to pogo element [Homo sapiens] R7942813 (39, 40) Salmonalla entertitiols] Novel Protein sim. GBank gil223050 prf 1817175B - nhaR gene [Salmonalla entertitiols] Novel Protein sim. GBank gil223050 prf 1817175B - nhaR gene [Salmonalla entertitiols] Novel Protein sim. GBank gil2493240 sp 010341 Y091_NPVOP - HYPOTHETICAL_29.9 KD PROTEIN (ORF92) Novel Protein sim. GBank gil2494533 sp P75939 F1GG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	394	100344205 (1889, 1890)		UNCLASSIFIED	1004, 1022, 1023,
101334754 (1343, 1344) Novel Protein sim. GBank gi[2144101 pir 155210 - tricarboxylate carrier - rat (fragment) 87940670 (547, 548) Novel Protein sim. GBank gi[2213611 (AC000103) - F21J9.5 100340738 (1613, 1614) Novel Protein sim. GBank gi[2225004 (U49973) - ORF1; MER37; putative transposase similar to pogo element [Homo sapiens] 87931075 (1577, 1578) Novel Protein sim. GBank gi[2225005 (U49973) - ORF2; function unknown [Homo sapiens] 87942813 (39, 40) Novel Protein sim. GBank gi[229050 pri 1817175B - nhaR gene [Salmonella enteritidis] Novel Protein sim. GBank gi[2494533 sp 070341 Y091_NPVOP - HYPOTHETICAL 29.3 KD PROTEIN (ORF92) 87940534 (529, 530) Novel Protein sim. GBank gi[2494533 sp 075939 FLGG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	395	100393146 (1679, 1680)	: gi 2129478 pir S51939 - chitinase (EC	UNCLASSIFIED	1022, 1033, 1040,
87940670 (547, 548) Novel Protein sim. GBank gi 2145684 pir S72805 - B1549_F2_87 protein - Mycobacterium leprae 100340738 (1613, 1614) Novel Protein sim. GBank gi 2213611 (AC000103) - F21J9.5 [Arabidopsis thallana] 87931075 (1577, 1578) Novel Protein sim. GBank gi 2226004 (U49973) - ORF1; MER37; putativo transposaes similar to pogo element [Homo sapiens] 87942813 (39, 40) Novel Protein sim. GBank gi 229050 prf 1817175B - nhaR gene [Salmonalla enteritidis] 87942813 (39, 40) Salmonalla enteritidis] 87942813 (1975, 1976) Novel Protein sim. GBank gi 2494533 sp P75939 FLGG_ECOLI - HYPOTHETICAL 29.3 KD PROTEIN (ORF92) Novel Protein sim. GBank gi 2494533 sp P75939 FLGG_ECOLI - FLAGEILAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	396	101334754 (1343, 1344)		UNCLASSIFIED	1014, 1022
100340738 (1613, 1614) Novel Protein sim. GBank gi[2213611 (AC000103) - F21J9.5 [Arabidopsis thaliana] 87931075 (1577, 1578) Novel Protein sim. GBank gi[2226004 (U49973) - ORF1; MER37; putative transposase similar to pogo element [Homo sapiens] 87938856 (1581, 1582) Novel Protein sim. GBank gi[2226005 (U49973) - ORF2: function unknown [Homo sapiens] 87942813 (39, 40) Novel Protein sim. GBank gi[229050 prf 1817175B - nhaR gene [Salmonella enteritidis] 87932519 (1975, 1976) Novel Protein sim. GBank gi[2493240 sp]O10341 Y091_NPVOP - HYPOTHETICAL 29.3 KD PROTEIN (ORF92) 87940534 (529, 530) Rovel Protein sim. GBank gi[2494533 sp]P75939 FLGG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	397	87940670 (547, 548)		UNCLASSIFIED	1022
87931075 (1577, 1578) Novel Protein sim. GBank gi[2226004 (U49973) - ORF1; MER37; putative transposase similar to pogo element [Homo sapiens] 87938856 (1581, 1582) Novel Protein sim. GBank gi[2226005 (U49973) - ORF2: function unknown [Homo sapiens] 87942813 (39, 40) [Salmonella enteritidis] 87942813 (39, 40) [Salmonella enteritidis] 87932519 (1975, 1976) Novel Protein sim. GBank gi[2494533 sp[P75939 FLGG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	398	100340738 (1613, 1614)		UNCLASSIFIED	1022
87938856 (1581, 1582) Novel Protein sim. GBank gi[2226005 (U49973) - ORF2: function unknown [Homo sapiens] 87942813 (39, 40) Novel Protein sim. GBank gi[229050]prf[]1817175B - nhaR gene [Salmonella enteritidis] 87942813 (1975, 1976) Novel Protein sim. GBank gi[2493240]sp]O10341 Y091 _ NPVOP - HYPOTHETICAL 29,3 KD PROTEIN (ORF92) Novel Protein sim. GBank gi[2494533]sp[P75939]FLGG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	399	87931075 (1577, 1578)		UNCLASSIFIED	1022
87942813 (39, 40) Novel Protein sim. GBank gi 229050 prf 1817175B - nhaR gene [Salmonella enteritidis] 87932519 (1975, 1976) Novel Protein sim. GBank gi 2493240 sp 010341 Y091_NPVOP - HYPOTHETICAL 29.3 KD PROTEIN (ORF92) 87940534 (529, 530) FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	400	87938856 (1581, 1582)		UNCLASSIFIED	1022
87932519 (1975, 1976) Novel Protein sim. GBank gi 2493240 sp O10341 Y091_NPVOP - HYPOTHETICAL 29,3 KD PROTEIN (ORF92) 87940534 (529, 530) Novel Protein sim. GBank gi 2494533 sp P75939 FLGG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	401	87942813 (39, 40)	unk gi 229050 prf 1817175B - nhaR gene	UNCLASSIFIED	1022
87940534 (529, 530) Novel Protein sim. GBank gi 2494533 sp P75939 FLGG_ECOLI - FLAGELLAR BASAL-BODY ROD PROTEIN FLGG (DISTAL ROD	402	87932519 (1975, 1976)		UNOLASSIFIED	1022
	403	87940534 (529, 530)		UNCLASSIFIED	1022

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